

# *In Vitro* Evaluation of the Antimethanogenic Potency and Effects on Fermentation of Individual and Combinations of Marine Macroalgae

Robert D. Kinley<sup>1</sup>, Matthew J. Vucko<sup>2</sup>, Loreнна Machado<sup>2</sup>, Nigel W. Tomkins<sup>1</sup>

<sup>1</sup>CSIRO Agriculture, Australian Tropical Science & Innovation Precinct, James Cook University, Townsville, Australia

<sup>2</sup>MACRO—The Centre for Macroalgal Resources and Biotechnology, College of Marine and Environmental Sciences, James Cook University, Townsville, Australia

Email: rob.kinley@csiro.au

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## Abstract

Contribution of ruminants to total greenhouse gas emissions in Australia is approximately 10% and likely to increase with demand for livestock products, thus an efficient method of mitigation must be implemented. The red marine macroalgae *Asparagopsis taxiformis* reduces enteric methane production by up to 99% *in vitro*. Other macroalgae with less potent antimethanogenic properties may complement inclusion of *Asparagopsis* in livestock feeds. Adoption of environmental based changes in livestock systems must provide benefits to producers if change in management is to be adopted. This study used 72 h *in vitro* fermentations with rumen inoculum to characterize and rank seven species of macroalgae at low inclusion that previously demonstrated some degree of antimethanogenesis at higher inclusion concentration. The seven were assessed at 5% inclusion (OM basis) and in combination with *Asparagopsis* to evaluate beneficial effects on fermentation. When tested individually, improvements in volatile fatty acids were generally observed, however, minimal effect on gas production and no clear justification for a ranking order were demonstrated. When tested in combination with *Asparagopsis*, the effects on fermentation were dominated by presence of *Asparagopsis* at 2% and no further benefits demonstrated. Therefore, *Asparagopsis* remains the only macroalga inducing near elimination of methane *in vitro* and benefit of combinations with other macroalgae evaluated in this study was not demonstrated. However, combination with high protein macroalgae is proposed to provide productivity enhancement during seasonal lows in grass quality and thus reduce methane emissions intensity providing a stronger conduit for environmental responsibility while increasing productivity.

## Keywords

*Asparagopsis*, Methane, Rumen, Seaweed, Algae

## 1. Introduction

A large proportion of methane ( $\text{CH}_4$ ) emitted into the atmosphere derives from agriculture, and specifically ruminant enteric fermentation which contributes approximately 28% of global anthropogenic  $\text{CH}_4$  emissions [1].

Ruminants rely on a complex rumen microbial consortium of bacteria, protozoa, archaea, fungi, and bacteriophages [2] associated with fermentation of feed. Fermentation of fibrous and nitrogenous feedstuff results in the production of volatile fatty acids (VFA) and microbial protein, used by the animal for growth, metabolism, and productivity [3]. During fermentation, carbon dioxide ( $\text{CO}_2$ ) and hydrogen ( $\text{H}_2$ ) are utilized by methanogenic archaea in the reductive methanogenesis pathway, which reduces partial  $\text{H}_2$  pressure, but also results in  $\text{CH}_4$  being emitted into the atmosphere [3]. Low  $\text{CH}_4$  producing ruminants tend to be more productive and increasing productivity of animals could also lessen  $\text{CH}_4$  emissions [4]. As a result, many strategies are being evaluated to mitigate enteric methanogenesis, including vaccines, bacteriocins and probiotics, bacteriophage therapy, genetic selection, feeding management and feed additives, and plant secondary metabolites [4].

Feed management and additives such as high-quality forages, grains, ionophores, fats, yeasts, enzymes, microbes, plant extracts and algae have the potential for  $\text{CH}_4$  abatement [5]. Algae products can improve ruminant health and productivity [6] [7], increase feed quality [8], and inhibit methanogenesis [9]. However, reducing enteric methanogenesis is challenging, and to be adopted as a methodology it needs to be sustainable, practical, economically viable, and improve animal productivity [4]. Recently, the marine macroalgae *Asparagopsis taxiformis* ([Delile] Trevisan de Saint-Léon 1845; *Asparagopsis*) has demonstrated effective inhibition of  $\text{CH}_4$  production *in vitro* at a dose of 2% of substrate organic matter (OM). In addition, *Asparagopsis* maintains the apparent *in vitro* degradability of OM (IVD-OM), and increases propionate but with a concomitant decrease of acetate [10] [11]. These rumen fermentation parameters are important indicators of fermentation efficiency [12]. Some other marine macroalgae that also reduce  $\text{CH}_4$ , but to a lesser extent than *Asparagopsis*, have demonstrated better anaerobic fermentation *in vitro* compared to equal supplementation (17%) with cotton seed meal by increasing the concentration of total VFA (TVFA) when included with a low-quality grass substrate [13]. Therefore, it may be possible to improve rumen fermentation using combinations of macroalgae at practical dose concentrations when paired with *Asparagopsis* by increasing VFA production and improving the VFA profile.

It was hypothesised that macroalgae demonstrating  $\text{CH}_4$  abatement *in vitro* at high dose concentrations would maintain antimethanogenesis at variable potency at low dose concentrations, and that when combined with the highly potent *Asparagopsis* they would enhance rumen fermentation. The aim of Experiment 1 was to rank seven select macroalgae for their potential as antimethanogenic feed additives using dose concentrations practical for livestock feeding, and establish their effects on fermentation parameters. The aim of Experiment 2 was to evaluate combining the seven macroalgae

with *Asparagopsis* to determine the effect of the combinations on methanogenesis and potential benefits to VFA production.

## 2. Materials and Methods

### 2.1. Selection and Preparation of Macroalgae and Rhodes Grass Substrates

The macroalgae species were selected to represent the major groups of marine macroalgae (red, brown, green) based on their demonstrated ability to decrease enteric CH<sub>4</sub> and improve fermentation [13] and are listed in **Table 1**. The biomass was sourced from either large scale culture at the Centre for Macroalgal Resources and Biotechnology (MACRO) James Cook University (JCU) in Townsville QLD (19.33°S; 146.76°E), Pacific Reef Fisheries in Ayr QLD (PRF; 19.58°S, 147.40°E), or intertidal flats near Townsville Qld at Nelly Bay (NB; 19°16'S; 146°85'E) with GBRMPA permit GO2/20234.1, or Rowes Bay (RB; 19°23'S; 146°79'E) with DPIF permit 103,256. The individual algae and sources were *Caulerpa taxifolia* ([M.Vahl] Agardh 1817; NB), *Cladophora patentiramea* ([Montagne] Kützing 1849; PRF), *Cystoseria trinodis* ([Forsskål] Agardh 1820; NB), *Dictyota bartayresii* (Lamouroux 1809; RB), *Padina australis* (Hauck 1887; RB), *Sargassum flavicans* ([Mertens] Agardh 1820; NB) and *Ulva ohnoi* (Hiraoka and Shimada 2004; MACRO).

Macroalgae were rinsed and centrifuged in a commercial washing machine at 1000 rpm for 5 min at ambient temperature to remove excess water. The biomass was freeze-dried (VirTis Benchtop 2K; Warminster PA, USA) at -55°C and 120 µbar for minimum 48 h, then milled to pass a 1 mm sieve and stored at -20°C. The basal substrate was Rhodes grass (*Chloris gayana*) with neutral detergent fibre (NDF) of 645 g·kg<sup>-1</sup> dry matter (DM) and acid detergent fibre (ADF) of 315 g·kg<sup>-1</sup> DM. The DM content of the various biomass was determined by drying at 105°C to constant weight and the OM content was determined by the combustion for 8 h at 550°C [14]. Crude protein and

**Table 1.** Compositional parameters of the macroalgae and Rhodes grass hay.

Substrate	Macroalgae Type	DM	OM	CP	GE <sup>a</sup>
<i>Asparagopsis taxiformis</i>	Rhodophyta (Red)	923 <sup>b</sup>	797	278	17
<i>Caulerpa taxifolia</i>	Chlorophyta (Green)	929	663	163	13
<i>Cladophora patentiramea</i>	Chlorophyta (Green)	939	564	120	11
<i>Cystoseria trinodis</i>	Ochrophyta (Brown)	909	650	92	12
<i>Dictyota bartayresii</i>	Ochrophyta (Brown)	940	658	90	13
<i>Padina australis</i>	Ochrophyta (Brown)	931	532	55	9
<i>Sargassum flavicans</i>	Ochrophyta (Brown)	919	646	42	12
<i>Ulva ohnoi</i>	Chlorophyta (Green)	898	695	215	12
Rhodes grass	-	916	804	167	17

a. MJ·kg<sup>-1</sup> DM. b. Parameters presented as g·kg<sup>-1</sup> unless otherwise stated. DM, dry matter; OM, organic matter; CP, crude protein.

gross energy quantification were determined as previously described by [13]. Crude protein of the macroalgae was based on total nitrogen (wt %) content using a nitrogen factor of 5 for the macroalgae [15] and 6.25 for the Rhodes grass.

## 2.2. Research Animals and Preparation of Rumen Fluid Inoculum

Rumen fluid inoculum (RF) was collected from four fistulated Brahman steers (*Bos indicus*; LW 460 ± 20 kg) fitted with 10 cm Bar Diamond (Parma, OH, USA) rumen cannulas. The steers were maintained at the College of Public Health, Medical and Veterinary Sciences of JCU according to current guidelines [16] and approved by the CSIRO animal ethics committee (A5/2011). The steers were maintained on Rhodes grass hay *ad libitum* for 6 months before the collection of RF, which was extracted 2 h after morning feeding by sampling from four quadrants of the rumen and hand squeezing to completely fill pre-warmed 1 L stainless steel thermos flasks.

## 2.3. Inoculation of *in Vitro* Fermentations

The RF was pooled and immediately processed by filtration through a 0.5-mm sieve and combined with incubation buffer (GVB) [17] at a ratio of 1:4 (RF:GVB). Throughout the inoculation process the RF buffer fermentation media (RFB) was maintained at 39°C and continuously mixed at 30% - 35% of maximum vortex speed (Major Science SWB 20 L<sup>-3</sup>; Saratoga, CA, USA) to ensure RFB homogeneity between individual fermentations. Anaerobic conditions were maintained with high purity N<sub>2</sub> (HPN<sub>2</sub>; BOC, Wetherill Park NSW, AUS). A Dose-It pump (Integra Biosciences, Hudson NH, USA) was used to aspirate 125 mL of RFB into each incubation bottle containing the macroalgae and Rhodes grass substrates. The substrate and macroalgae were preloaded prior to RF collection into 250 mL Simax incubation bottles (Sázava, CZE). Each bottle was purged HPN<sub>2</sub>, capped, and warmed to 39°C. The bottles were sealed with an Ankom RF1 gas production module (Macedon NY, USA), placed in mixing incubators (Ratek OM11; Boronia VIC, AUS), and maintained at 39°C and oscillating at 85 RPM for 72 h.

## 2.4. Experimental Design

Experiment 1 was conducted to rank macroalgae according to their antimethanogenic potency and effects on rumen fermentation parameters. This was accomplished using *in vitro* fermentation batch culture (Ankom) to determine effects of macroalgae inclusion on total gas (TGP), CH<sub>4</sub>, and VFA production, and IVD-OM. A series of four incubation periods were completed and consisted of 1.0 g of Rhodes grass substrate as control and appropriate quantities of each macroalga to achieve a dose concentration of 5% of the substrate OM according to the biomass composition described in **Table 1**. Fermentations were also completed as controls (no macroalgae), a positive-control (P-control) of *Asparagopsis* at 2%, and RFB blanks. The inclusion of the macroalgae at 5% was determined as an arbitrary feasible level of feeding for livestock and based on results previously described for inclusion approaching 20% [13]. The inclusion of *Asparagopsis* at 2% was set based on previously determined optimum dose for this sea-

weed in rumen fermentations [10] [11]. The seven macroalgae and *Asparagopsis* were randomly assigned to the four incubation periods ( $n = 3$ ), and controls and blanks were included in all periods ( $n = 4$ ). Within each incubation period, there was duplication of each macroalga, and controls and blanks at each sampling time point (12, 24, 48, and 72 h).

Experiment 2 was conducted to determine if the seven macroalga when combined with *Asparagopsis* studied in Experiment 1 would benefit *in vitro* rumen fermentation. This was accomplished using the same fermentation techniques as Experiment 1. A series of five incubation periods were completed and consisted of 1.0 g of Rhodes grass substrate as control and appropriate quantities of each macroalga to achieve a dose concentration of 5% (OM basis) in combination with *Asparagopsis* at 2% according to the biomass composition described in **Table 1**. Fermentations were also completed as controls (no macroalgae), a P-control of *Asparagopsis* at 2%, and RFB blanks. The seven combinations and *Asparagopsis* were randomly assigned to the five periods ( $n = 3$ ), and controls and blanks were included in all periods ( $n = 5$ ). Within each incubation period, there was duplication of each macroalga combination, and controls and blanks at each sampling time point (12, 24, 48, and 72 h).

## 2.5. Total Gas and Methane Production

The fermentation methods used in this study were similar to [18] but using the Ankom RF1 system and gas analysis as described by [10]. Pressure accumulation in the incubation bottles was measured continuously and recorded every 20 min. The maximum fermentation pressure inside the incubation bottle was set to 3 psi which, when exceeded, caused venting for 250 ms and the pressure change accounted in the cumulative pressure recording. Gas pressure was measured every 60 s and cumulative pressure was recorded at 20 min intervals. The TGP was expressed as mL of gas produced per gram of substrate OM ( $\text{mL}\cdot\text{g}^{-1}$  OM) by application of the natural gas law to the accumulation of the recorded gas pressure while accounting for individual bottle volume.

*In vitro*  $\text{CH}_4$  production was quantified in time series at multiple time points of 12, 24, 48 and 72 h during the fermentation. The mL of  $\text{CH}_4$   $\text{g}^{-1}$  of substrate OM was estimated using concentrations in headspace at the time series points while assuming homogeneity of headspace gas. The headspace samples were collected into 10 mL Labco Exetainer vials (Lampeter, GBR) and quantified using gas chromatography (GC) according to parameters described by [10]. Concentrations of  $\text{CH}_4$  in headspace gas were converted to  $\text{mL}\cdot\text{g}^{-1}$  substrate OM based on TGP at the relative time series points and assuming headspace homogeneity at system venting [19].

## 2.6. Apparent *in Vitro* Digestible Organic Matter and Volatile Fatty Acids

The IVD-OM and VFA production was determined as described by [10] and quantified to coincide with  $\text{CH}_4$  determinations at each time series point. Each fermentation was chilled to cease bacterial activity and the *in vitro* fluid (IVF) was then vacuum filtered through a Duran No. 1 porosity glass fritted crucible containing a 0.5 cm layer of sand

filtration aid. The pH was measured on the filtrate. Crucibles and wet residue were oven-dried to constant weight at 105°C for DM determination. Residue OM was determined as loss on ignition in a muffle furnace at 550°C for 8 h [14].

The TVFA accumulated in the IVF were quantified after termination of fermentation. Preparation of IVF prior to GC analysis consisted of addition 4 mL of IVF to 1 mL of 20% metaphosphoric acid containing 11 mM of 4-methylvaleric acid (Sigma-Aldrich; Castle Hill NSW, AUS) providing 2.2 mM internal standard. The samples were mixed and stored at -20°C until a 1.5 mL subsample was centrifuged for 15 min at 13,500 rpm and 4°C (Labnet Prism R; Edison NJ, USA). The supernatant was filtered through 0.2 µm PTFE syringe tip filters (Agilent; Santa Clara, CA, USA) and quantified by GC according to parameters described by [10].

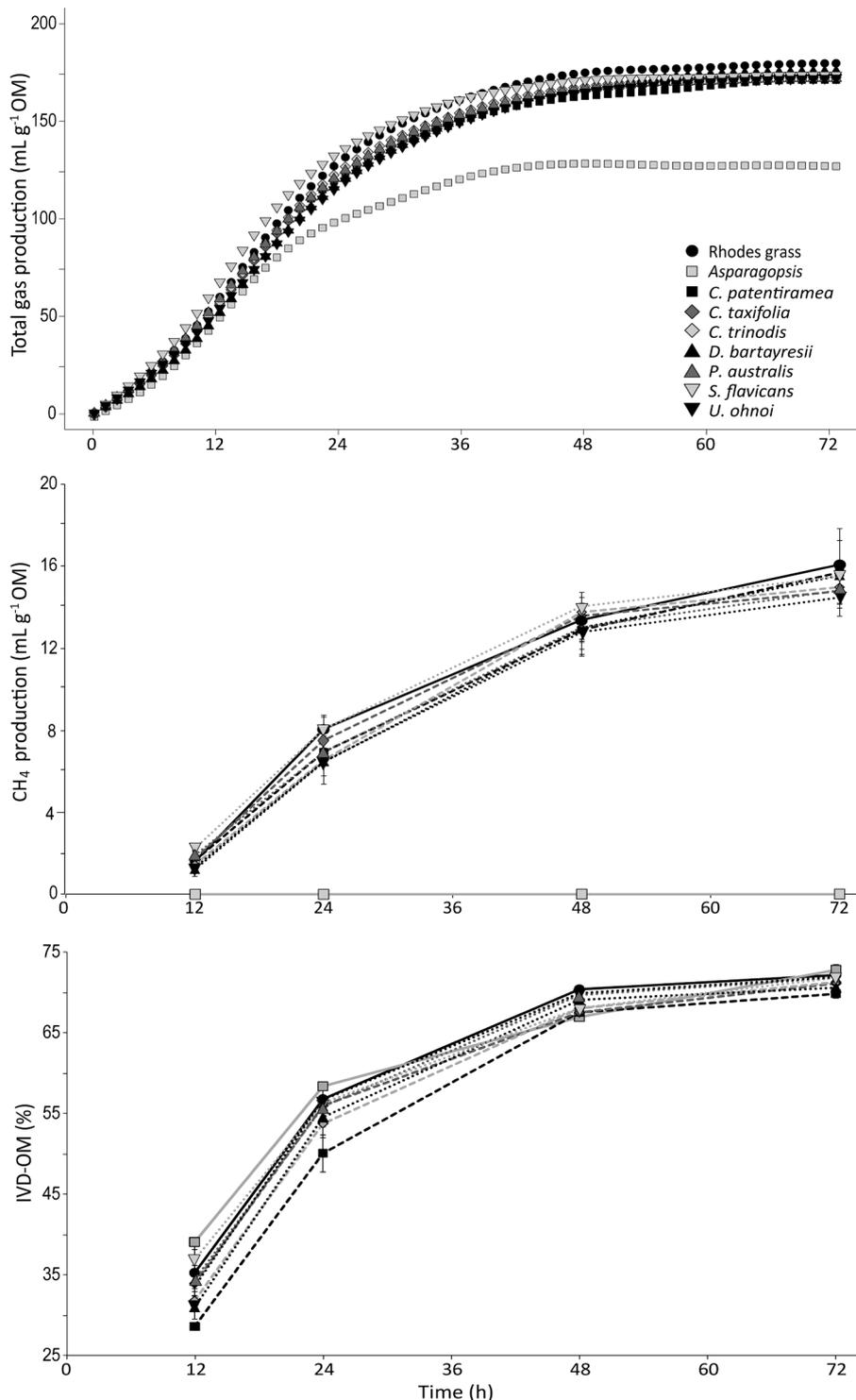
## 2.7. Statistical Analysis

Two-factor repeated-measures permutational analysis of variance (PERMANOVA) was used to test for significant differences in TGP, CH<sub>4</sub> production, and IVD-OM over time. A one-factor PERMANOVA was used to test for significant differences in the production of VFA between treatments (fixed factor) using Primer 6 (version 6.1.13; [20] statistical software and PERMANOVA+ (version 1.0.3) [21]. For PERMANOVA, Bray-Curtis similarity matrices were produced using the untransformed raw data and dummy variables (0.0001) were used to account for zero values. The *P*-values were calculated from 999 (TGP) and 9,999 (CH<sub>4</sub>, IVD-OM, and VFAs) random permutations. Pair-wise *a posteriori* comparison was used to determine significant groupings, where applicable. For PERMANOVA, differences were considered significant if *P* < 0.05. The TGP data were also fitted with generalized additive models (GAM) using cubic regression spline smoothers to predict the relationship and examine differences between TGP over time. The generalized additive models were produced using the mgcv package within the R language (version 3.0.1) [22]. Goodness-of-fit of the individual smoothers was quantified using the hydroGOF package within the R language and was assessed from the proportion of variance in the data that was accounted for by the model (*r*<sup>2</sup>) [23].

## 3. Results

### 3.1. Experiment 1: Ranking

Based on predictions of the GAM, which had a high goodness-of-fit (*r*<sup>2</sup> > 0.97), there was a significant difference (*P* < 0.001) in TGP in the fermentations over 72 h (**Figure 1**). In addition, the TGP reached its plateau approaching 48 h in all fermentations and the *Asparagopsis* induced a lower TGP rate compared with all other treatments (*P* < 0.001). There was no significant difference in cumulative TGP between the seven macroalgae with inclusion at 5% (OM basis) and the Rhodes grass control after 72 h of fermentation. The TGP for all macroalgae other than *Asparagopsis* ranged between 171 and 176 mL·g<sup>-1</sup> OM, representing an insignificant 2.4% - 4.8% reduction in TGP com-



**Figure 1.** The effect of inclusion of seven different macroalgae on gas production and substrate digestibility over 72 h of *in vitro* fermentation with rumen fluid. From top down: Total gas production (TGP); CH<sub>4</sub> production; and apparent *in vitro* digestibility (IVD-OM). The Rhodes grass control substrate was equal in all fermentations, the *Asparagopsis* control was included at a concentration of 2% of substrate OM, and the other seven macroalgae included at 5%. No ±SE is presented for TGP because SE was smaller than the symbols.

pared to the Rhodes grass control ( $180 \text{ mL}\cdot\text{g}^{-1} \text{ OM}$ ). However, the *Asparagopsis* at 2% had significantly less TGP ( $127 \text{ mL}\cdot\text{g}^{-1} \text{ OM}$ ) representing a 26% - 29% reduction compared to all other macroalgae and controls.

The  $\text{CH}_4$  production as affected by inclusion of the seven macroalgae other than *Asparagopsis* was not altered significantly compared to the Rhodes grass control during the course of the 72 h incubations (Figure 1). The rate of  $\text{CH}_4$  production as defined by time series headspace sampling was variable in the same manner as for TGP. The only significant reduction in  $\text{CH}_4$  was induced by *Asparagopsis* in the same way as previously demonstrated [10] and no detectable  $\text{CH}_4$  was produced. The  $\text{CH}_4$  production from the other fermentations ranged from a low of  $14 \text{ mL}\cdot\text{g}^{-1} \text{ OM}$  to a high of  $16 \text{ mL}\cdot\text{g}^{-1} \text{ OM}$  for *U. ohnoi* and the Rhodes grass control, respectively.

A typical pattern of IVD-OM increasing with time was demonstrated with or without macroalgae (Figure 1), however, the rate of digestion of the substrate varied between treatments in the first half of the fermentation period ( $P = 0.004$ ). *Asparagopsis* induced the earliest onset of substrate degradation in the same fashion as demonstrated by [10] for *Asparagopsis* at dose concentrations  $\leq 5\%$  of substrate OM. The *C. patentiramea* was the slowest to achieve maximum rate of IVD-OM. Approaching 48 h, IVD-OM in all the fermentations coalesced, and after 72 h had similar IVD-OM ranging from a low of 70% to a high of 72% for *C. patentiramea* and the *Asparagopsis*, respectively. In the absence of macroalgae the IVD-OM for the Rhodes grass control was 72%.

In Experiment 1 the series of *in vitro* fermentations of the seven macroalgae or *Asparagopsis* P-control did not induce significant differences between any of the treatments or Rhodes grass control for TVFA ( $P = 0.093$ ) or propionate ( $P = 0.098$ ). However, fermentations including the *Asparagopsis* were significantly lower for acetate ( $P = 0.002$ ) and higher for butyrate ( $P = 0.004$ ) compared to the other fermentations (Table 2). A significant increase in propionate and butyrate with 2% *Asparagopsis* inclusion has been previously reported [10], however this feature was slightly dampened in the present study for propionate but confirmed the previously observed butyrate concentrations. Compared to the Rhodes grass control, after 72 h of fermentation the macroalgae induced marginal reductions in TVFA on a molar concentration basis from a low of 9% to a high of 23% for *D. bartayresii* and *C. patentiramea*, respectively, and *Asparagopsis* reduced TVFA by 22%. The production of acetate was marginally reduced by a low of 10% to a high of 24% by inclusion of *D. bartayresii* and *C. patentiramea*, respectively, and a significant 44% with *Asparagopsis*, respectively. Conversely, propionate was marginally decreased by a low of 5% to a high of 24% with inclusion of *D. bartayresii* and *C. patentiramea*, respectively, and increased by 9% with *Asparagopsis*. The production of butyrate was marginally decreased by a low of 11% to a high of 37% for *D. bartayresii* and *C. patentiramea*, respectively, and significantly increased by 76% with *Asparagopsis*.

There were not adequate differences in TGP,  $\text{CH}_4$ , or VFA production to justify well defined ranking for the seven macroalgae evaluated. Clearly, none of the seven macro-

**Table 2.** The effect of inclusion of seven macroalgae species on accumulation of volatile fatty acids after 72 h of *in vitro* fermentation with rumen fluid. The Rhodes grass control substrate was equal in all fermentations, the *Asparagopsis* P-control was included at 2% of substrate OM, and the other seven macroalgae included at 5%.

Treatment	Volatile Fatty Acids (mM)			
	Total	Acetate	Propionate	Butyrate
Rhodes grass	43.87 ± 0.82	28.88 ± 0.41	11.62 ± 0.32	1.86 ± 0.06
<i>Asparagopsis taxiformis</i>	34.19 ± 0.50	16.28 ± 0.41 <sup>a</sup>	12.65 ± 0.10	3.27 ± 0.11 <sup>a</sup>
<i>Caulerpa taxifolia</i>	38.69 ± 0.92	25.01 ± 0.28	10.86 ± 0.57	1.51 ± 0.23
<i>Cladophora patentiramea</i>	33.81 ± 3.17	22.10 ± 2.09	9.46 ± 1.02	1.17 ± 0.17
<i>Cystoseria trinodis</i>	38.73 ± 3.07	24.85 ± 2.51	11.03 ± 0.44	1.58 ± 0.26
<i>Dictyota bartayresii</i>	40.04 ± 1.16	26.11 ± 1.21	11.05 ± 0.46	1.66 ± 0.23
<i>Padina australis</i>	37.24 ± 0.35	23.66 ± 0.57	10.71 ± 0.40	1.56 ± 0.20
<i>Sargassum flavicans</i>	37.32 ± 1.12	23.97 ± 0.62	10.64 ± 0.47	1.44 ± 0.28
<i>Ulva ohnoi</i>	37.01 ± 3.63	24.02 ± 2.72	10.22 ± 0.91	1.45 ± 0.14

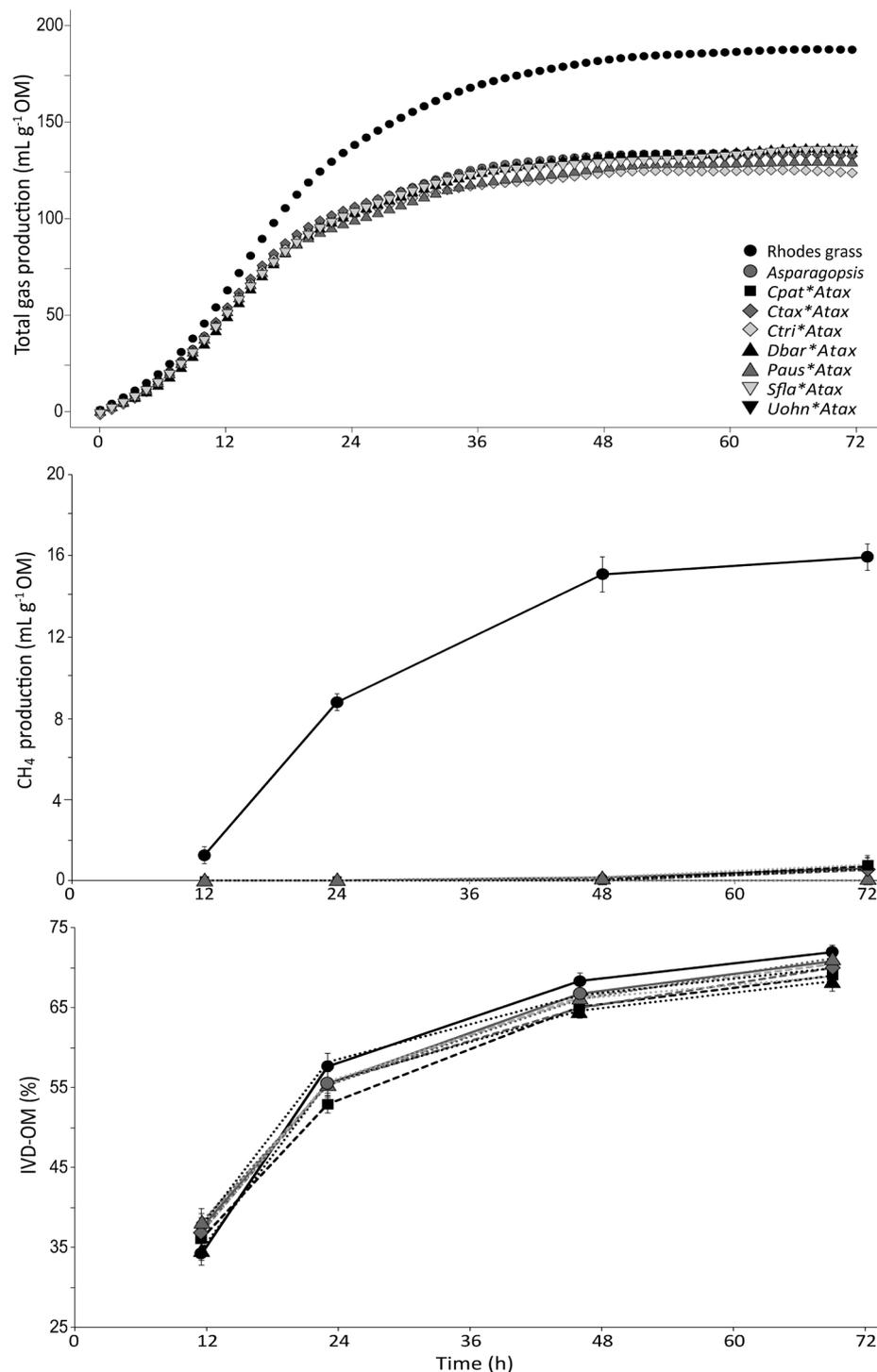
a. Significant at  $P < 0.05$ .

algae at 5% of substrate OM compares to *Asparagopsis* at 2% for CH<sub>4</sub> abatement without detriment to IVD-OM during *in vitro* fermentation. Therefore based solely on numerical CH<sub>4</sub> abatement the ranking order from lowest to highest CH<sub>4</sub> produced is as follows: 1) *U. ohnoi*; 2) *C. taxifolia*; 3) *P. australis*; 4) *C. trinodis*; 5) *D. bartayresii*; 6) *S. flavicans*; 7) *C. patentiramea*.

### 3.2. Experiment 2: Combinations

Similar to the results of Experiment 1, based on predictions of the GAM, which had a high goodness-of-fit ( $r^2 > 0.98$ ), there was a significant difference ( $P < 0.001$ ) in TGP in the fermentations over 72 h (Figure 2). In addition, the TGP reached its plateau approaching 48 h in all fermentations, however, in contrast with Experiment 1, the combination treatments were equivalent to the *Asparagopsis* P-control at 2% of substrate OM in the reduction of TGP, but all were significantly different from the Rhodes grass control ( $P < 0.001$ ) after 72 h of fermentation. The TGP for the combinations ranged between 124 and 136 mL·g<sup>-1</sup> OM representing a significant 28% - 34% reduction in TGP compared to the Rhodes grass control (188 mL·g<sup>-1</sup> OM). The *Asparagopsis* induced a TGP of 134 mL·g<sup>-1</sup> OM and therefore a 28% reduction, similar to the combinations demonstrating an overwhelming effect of *Asparagopsis* when combined with other macroalgae.

The CH<sub>4</sub> production as affected by inclusion of the seven macroalgae combined with *Asparagopsis* was not different compared to the *Asparagopsis* alone during the course of the 72 h incubations (Figure 2). In the absence of macroalgae the fermentation initiated production of CH<sub>4</sub> immediately after inoculation with RFB, however, the inclusion of the macroalgae combinations completely inhibited CH<sub>4</sub> until 48 h when trace



**Figure 2.** The effect of inclusion of seven different macroalgae combined with *Asparagopsis* on gas production and substrate digestibility over 72 h of *in vitro* fermentation with rumen fluid. From top down: Total gas production (TGP); CH<sub>4</sub> production; and apparent *in vitro* digestibility (IVD-OM). The Rhodes grass control substrate was equal in all fermentations, the *Asparagopsis* control was included at a concentration of 2% of substrate OM, and the seven macroalgae combinations resulted in an inclusion of *Asparagopsis* at 2% and each of the seven macroalgae at 5%. No  $\pm$ SE is presented for TGP because SE was smaller than the symbols.

amounts were measured for the *S. flavicans* combination (0.2 mL·g<sup>-1</sup> OM) representing 98% inhibition. After 48 h of fermentation *S. flavicans* and *D. bartayresii* combined with *Asparagopsis* accumulated small amounts of CH<sub>4</sub> such that at 72 h of fermentation these two combinations had reduced CH<sub>4</sub> by 95% (0.8 mL·g<sup>-1</sup> OM) compared to the Rhodes grass control (16.0 mL·g<sup>-1</sup> OM). The *Asparagopsis* alone reduced CH<sub>4</sub> by 99% (0.2 mL·g<sup>-1</sup> OM) after 48 h and by 96% (0.6 mL·g<sup>-1</sup> OM) at 72 h.

In the same way as Experiment 1 the typical pattern of IVD-OM increasing with time was demonstrated with or without macroalgae combinations (Figure 2). However, in Experiment 2 IVD-OM induced by the macroalgae combined with *Asparagopsis* was not as variable throughout the 72 h *in vitro* fermentations. Other than the Rhodes grass control, all fermentations contained 2% *Asparagopsis* which negated effects of the other macroalgae in the combinations. All the fermentations coalesced from onset of fermentation and after 72 h of fermentation the IVD-OM ranged from a low of 68% to a high of 72% for the *D. bartayresii* combined with *Asparagopsis* and Rhodes grass control, respectively.

In Experiment 2, *in vitro* fermentations with inclusion of the seven macroalgae combinations or *Asparagopsis* induced significant reductions in TVFA ( $P = 0.002$ ) and acetate ( $P < 0.001$ ), and significant increases in propionate ( $P < 0.001$ ) and butyrate ( $P < 0.001$ ) compared with the Rhodes grass control (Table 3). However, there was no significant difference between the *Asparagopsis* alone and the combinations, therefore improvements in VFA profiles due to combining *Asparagopsis* with any of the seven candidate macroalgae was not demonstrated. Compared with the Rhodes grass control, the macroalgae combinations induced reduction in TVFA on a molar concentration

**Table 3.** The effect of inclusion of seven macroalgae species combined with *Asparagopsis* on accumulation of volatile fatty acids after 72 h of *in vitro* fermentation with rumen fluid. The Rhodes grass control substrate was equal in all fermentations, the *Asparagopsis* P-control was included at 2% of substrate OM, and the seven macroalgae combinations resulted in inclusions of *Asparagopsis* at 2% and each of the seven macroalgae at 5%.

Treatment	Volatile Fatty Acids (mM)			
	Total	Acetate	Propionate	Butyrate
Rhodes grass	43.32 ± 0.12 <sup>a</sup>	31.56 ± 0.14 <sup>a</sup>	7.88 ± 0.12 <sup>a</sup>	2.11 ± 0.06 <sup>a</sup>
<i>Asparagopsis taxiformis</i>	37.26 ± 1.10	21.83 ± 0.62	9.60 ± 0.19	3.49 ± 0.28
<i>Caulerpa taxifolia</i> ‡	40.29 ± 0.87	23.97 ± 0.74	9.98 ± 0.10	3.88 ± 0.06
<i>Cladophora patentiramea</i>	39.32 ± 0.65	23.38 ± 0.33	9.84 ± 0.13	3.76 ± 0.24
<i>Cystoseria trinodis</i>	38.77 ± 1.16	22.71 ± 0.68	9.91 ± 0.36	3.79 ± 0.13
<i>Dictyota bartayresii</i>	38.64 ± 0.53	23.09 ± 0.27	9.79 ± 0.11	3.54 ± 0.21
<i>Padina australis</i>	38.14 ± 0.05	22.69 ± 0.24	9.63 ± 0.18	3.57 ± 0.18
<i>Sargassum flavicans</i>	38.23 ± 1.22	22.46 ± 1.10	9.82 ± 0.17	3.58 ± 0.15
<i>Ulva ohnoi</i>	39.10 ± 0.77	23.52 ± 0.85	9.72 ± 0.10	3.54 ± 0.21

a. Significant at  $P < 0.05$ . ‡Each of the seven macroalgae represents half of a combination pair with.

basis from a low of 7% to a high of 12% and 14% by inclusion of the *P. australis* and *C. taxifolia* combined with *Asparagopsis* and *Asparagopsis* alone, respectively. The production of acetate was reduced by a low of 24% to a high of 29% and 31% by inclusion of *S. flavicans* and *C. taxifolia* combined with *Asparagopsis*, and *Asparagopsis* alone, respectively. Conversely, propionate was increased by a low of 21% to a high of 26% and 21% by inclusion of the *P. australis* and *C. taxifolia* combined with *Asparagopsis*, and *Asparagopsis* alone, respectively. The production of butyrate was increased by a low of 75% to a high of 91% and 72% for the *U. ohnoi* and *C. taxifolia* combined with *Asparagopsis*, and *Asparagopsis* alone, respectively.

#### 4. Discussion

This report represents the only current study of these seven macroalgae species (**Table 1**) at low inclusion *in vitro*. The purpose of this study was to rank them for beneficial effects *in vitro* prior to selection of one candidate for evaluation *in vivo*. However, there was no clear ranking order based on CH<sub>4</sub> mitigation and improvements in *in vitro* rumen fermentation. In combining the seven macroalgae with *Asparagopsis* there was no significant difference between the combinations and in all cases the *Asparagopsis* overwhelmed the effects on fermentation which eclipsed potential benefits of combining these macroalgae.

These macroalgae have been evaluated in rumen fermentations *in vitro* at high dose approaching 20% of substrate OM and the effects on TGP was variable and mostly significant reductions compared to controls were reported [13] [24]. In those studies the reductions in TGP were concomitant with reductions in CH<sub>4</sub> and in some cases detrimental effect on IVD-OM. Fermentations with the individual (pure) macroalgae as inclusions with Rhodes grass in Experiment 1 of the present study also demonstrated variable TGP. However, lack of significant reduction can be attributed to the much lower inclusion concentration of 5% in the fermentations. It was hypothesized that the effect demonstrated at high dose would remain at 5%, however only *Asparagopsis* at 2% maintained its potency at low dose which was a direct result of methanogenesis inhibition. The halogenated bioactive compounds inherent in *Asparagopsis* spp. have potent antimethanogenic properties [25] that are not inherent in the other macroalgae evaluated in the present study. Their mode of action at high dose may be attributed to organic acids, tannins, phlorotannins, polyphenolics, aminoglycans and other compounds that have antibacterial or inhibitory effect on rumen microbial metabolism, but have minimal effect at a low dose.

The reduction in CH<sub>4</sub> induced by these macroalgae has been reproduced at high inclusion concentrations *in vitro* [13] [24], unfortunately in the same way as TGP, at 5% inclusion, the effect was minimal. However, the *Asparagopsis* at 2% was consistent with previous studies specific to *that* macroalga demonstrating nearly complete elimination of CH<sub>4</sub> production *in vitro* [10] [11]. Without significant differences in TGP and CH<sub>4</sub> the ranking order was not clear and only numerical differences could be applied in the ranking. Thus, currently the *Asparagopsis* spp. are the only macroalgae demonstrating

CH<sub>4</sub> abatement ability exceeding 70% reduction at low dose and typically abatement > 99% is demonstrated.

Early in the fermentations the IVD-OM was variable with some of the macroalgae inducing a lag in the onset of fermentation. Previous research has shown a lag in TGP with addition at high dose [24] and this feature appears to be present at low dose, however both TGP and IVD-OM coalesced approaching 48 h of fermentation, and except for TGP induced by *Asparagopsis*, all fermentations were equal after 72 h. The *Asparagopsis* did not have different effect compared to the other macroalgae on IVD-OM demonstrating the importance of dose concentrations of macroalgae on rumen fermentation *in vitro* and presumably in ruminant animals. This indicates the importance of low dietary concentrations to maintain rumen efficiency and the seven macroalgae used in Experiment 1 demonstrated little benefit on IVD-OM or CH<sub>4</sub> emissions at low dose. An important feature to note is that none of the macroalgae had a negative effect on IVD-OM at the dose concentration studied.

The effect of inclusion of antimethanogenic compounds on production of VFA during *in vitro* fermentations has been variable in most studies, however, a trend concomitant with significant CH<sub>4</sub> reduction is in favour of increased propionate [26]. This has also been reported in previous studies using macroalgae which sometimes demonstrate a decrease in acetate in favour increased propionate and is more prominent with increasing dose. Macroalgae species that have a moderate or weak antimethanogenic capacity may not induce changes in the VFA profile [11] [27]. In the present study it was apparent that the seven macroalgae are weak antimethanogenic agents *in vitro* and as such they have minimal effects on VFA production. The TVFA and acetate results at 5% inclusion indicate a variable and small but not significant reduction with no apparent change in propionate or butyrate. These same macroalgae induced an increase in propionate with decreasing acetate at high dose, however, at high dose some species decreased TVFA as a result of detriment to IVD-OM [13]. It is important to maintain or improve IVD-OM to maximize fermentation efficiency which reiterates the requirement for appropriate dose concentrations of any dietary inclusion. Notably, Experiment 1 demonstrated that six of the seven macroalgae (excluding *C. patentiramea*) induced production of marginally more TVFA and significantly more acetate than *Asparagopsis* and so demonstrated their potential to enhance fermentation efficiency when included in combination with *Asparagopsis*.

It was hypothesized that combining macroalgae with *Asparagopsis in vitro* would improve fermentation because these were identified in Experiment 1 as weakly antimethanogenic and resulted in some increase in VFA compared with *Asparagopsis* alone. However, at 5% in combination with *Asparagopsis* at 2% there was no evidence demonstrating improvements at a level to provide incentive for a follow-up *in vivo* study.

There was some variable but small decreases in TGP for some of the combinations compared with *Asparagopsis* alone, however all were significantly reduced compared to fermentations without macroalgae. The universal decrease in TGP was attributed to the

*Asparagopsis* proportion of the combinations and a direct result of the near elimination of CH<sub>4</sub> production. However, a minimal concentration of CH<sub>4</sub> was detected in the final measurement (72 h) suggesting a gradual loss in antimethanogenic ability over time at the dose concentration used in this study. This phenomenon was described with *Asparagopsis* at doses ≤2% of substrate OM [10]. In the present study this occurred with *S. flavicans* and *D. bartayresii* combined with *Asparagopsis*, however, this does not indicate an effect specific to these combinations as a minimal level of CH<sub>4</sub> was detected after 72 h with the *Asparagopsis* alone. Total depletion of CH<sub>4</sub> is not expected *in vivo* where rumen fermentation is much more robust than in batch cultures. Also, feed residence time in the rumen is typically less than 72 h [28].

The loss of energy as CH<sub>4</sub> has potential to be reclaimed as productivity, however this can't be demonstrated *in vitro* and *in vivo* studies are necessary to determine the extent of productivity gains. When demonstrated *in vivo* productivity gains would dramatically increase the value of *Asparagopsis* and macroalgae combinations for livestock production systems. Adoption of any CH<sub>4</sub> mitigation strategy requires more than environmental benefits. The value of carbon abatement may eventually provide revenue incentive for producers to adopt macroalgae feed additives based on CH<sub>4</sub> abatement. However, improvements in productivity enhance the environmental value of macroalgae.

The IVD-OM was not negatively affected, however, the hypothesis of improvements to fermentation was not demonstrated by the combinations compared to *Asparagopsis*. All the fermentations, with and without macroalgae were stable and IVD-OM was not different. Improvements in IVD-OM would provide a conduit for improved utilization of feed and offset the cost of supplementation with macroalgae. Further investigation into macroalgae on improved feed energy utilization, productivity, and feed quality is necessary, particularly relative to periods of poor grass quality for grazing livestock [29].

In light of the VFA results of previous work [13] and Experiment 1 it was hypothesized that when combined with *Asparagopsis* the other macroalgae would have improved VFA production compared to the *Asparagopsis* alone. This effect unfortunately was not demonstrated and the combinations adopted similar profiles as *Asparagopsis* alone and the variability between treatments observed in Experiment 1 was muted in Experiment 2, thus again demonstrating the dominant effect of *Asparagopsis in vitro*. It is typical with *Asparagopsis* that decrease in CH<sub>4</sub> is concomitant with decrease in acetate and increase in propionate. It is common for antimethanogenic feed additives to have this effect *in vitro* [10] and *in vivo* [30] [31] and is believed to be due to reductive propionate production being more favourable than acetogenesis in the presence of excess hydrogen [26].

Although the present study did not support the use of macroalgae combinations to decrease CH<sub>4</sub> production *in vitro* the utility of combinations of macroalgae to enhance ruminant animal productivity and reduce CH<sub>4</sub> emissions is worthy of further exploration. Supplementation of high protein macroalgae such as the freshwater green *Oe-*

*dogonium* sp. is feasible up to 25% of intake [11] which could increase the proportion of rumen bypass protein thus benefiting productivity [32]. Alternative sources of protein can also reduce CH<sub>4</sub> emissions intensity by improving productivity of grass fed beef during those periods of decreasing diet quality. Supplementation with macroalgae can therefore directly reduce methanogenesis and reduce emission intensity by improving the product to emissions ratio.

## 5. Conclusion

There was not enough difference induced in rumen fermentation efficiency or CH<sub>4</sub> production *in vitro* to support a conclusive ranking order between the seven individual macroalgae at the 5% inclusion concentration of this study. A numerical difference indicates *U. ohnoi* and *C. patentiramea* were the most and least antimethanogenic, respectively. When macroalgae were combined with *Asparagopsis*, a known potent antimethanogenic agent *in vitro*, there was not an adequate effect to justify proceeding to *in vivo* evaluation or recommendation for use of the combinations in livestock feed. However, high protein macroalgae supplemented at higher dietary concentrations may provide greater benefit when combined with *Asparagopsis* by contributing to reduced CH<sub>4</sub> emissions through further improved productivity at times of low feed quality thus reducing emissions intensity per product output.

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